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Allelic forms of rat kappa chain genes: evidence selection at the level of nucleotide sequence.";
Proc. Natl. Acad. Sci. U.S.A. 78:7064-7068(1981).
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Milelic forms of rat kappa chain genes: evidence selection at the level of nucleotide sequence.";
Proc. Natl. Acad. Sci. U.S.A. 78:7064-7068(1981).
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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
IG KAPPA CHAIN C REGION, B ALLELE.
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PIR; A02117; K1RTB.
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MEDLINE-75212238; PubMed-807630;
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FRAMEWORK 2.
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FRAMEWORK 3.
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MEDLINE-83271467; PubMed-6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein Markham A.F., Milstein Markham Sequences define an unusually restricted IGG remarks.
                                                 MEDLINE-83271467; PubMed-6877353; Kaartinen M., Griffiths G.M., Markham A.F., Milstein "mRNA sequences define an unusually restricted IgG rephenyloxazolone and its early diversification."; Nature 304.320-324(1983).
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-!- FUNCTION: ANTI-2-PHENYL OXAZOLONE (PHOX) ANTIBODY.
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Mammalia; Eutheria;
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13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG KAPPA CHAIN V-VI REGION NQ5-61.1.2.
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arvota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Immunoglobulin V region; Hybridoma.
Immunoglobulin V region; Hybridoma.
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                                                                                                                                                                                                                                                                                                                                                             Nature 304:320-324(1983).
-!- FUNCTION: ANTI-2-PHENYL OXAZOLONE (PHOX) ANTIBODY.
                                                                                                                                                                                                                                                                                                                                                                                 "mRNA sequences define an unusually restricted IgG
phenyloxazolone and its early diversification.";
                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=83271467;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                   the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
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15-JUL-1999
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FSGSGSGTSYSLTISSVEAEDAATYYCQQYSGY-RTFGGGTKLEIKR 108
                   VLTQSPAIMSASPGEKVTMTCRASSSVSSSYLHWYRQKSGASPKLWIYSTSNLASGVPAR 62
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Pro; IPR003006; -.
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                                                                       Similarity
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11556 MW;
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11605 MW;
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                                                             Score 421.5; DB
Pred. No. 1e-26;
9; Mismatches
                                                                                                                                    COMPLEMENTARITY-DETERMINING FRAMEWORK 4.
BY SIMILARITY.
                                                                                                                                                                    FRAMEWORK
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BY SIMILARITY
                                                                                                                  72488DA9EF354934 CRC64;
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Best Local :
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13-AUG-1987
13-AUG-1987
15-JUL-1999
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SEQUENCE
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                                                                      KV6A_MOUSE STANDARD; PRT; 107 AA. P01675; 21-JUL-1986 (Rel. 01, Created) 21-JUL-1986 (Rel. 01, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) 15 KAPPA CHAIN V-VI REGION XRPC 44.
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MEDLINE-83271467; PubMed-6877353;
Kaartinen M., Gritfiths G.M., Markham A.F., Mil
"MRNA sequences define an unusually restricted."
                                      Mus musculus (Mouse)
Eukaryota; Metazoa; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
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Pfam; PF00047; ig; 1
Immunoglobulin V reg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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-!- FUNCTION: ANTI-2-PHENYL OXAZOLONE (PHOX) ANTIBODY.
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-AUG-1987 (Rel. 05, Last sequence update)
-JUL-1999 (Rel. 38, Last annotation updat
KAPPA CHAIN V-VI REGION NQ5-78.2.6.
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                   Rodentia;
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Pred. No. 1.2e
8; Mismatches
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BY SIMILARITY.
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                   Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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RESULT 11
KV6D_MOUSE
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Best Local :
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HSSP; P01679; 2FBJ.
HSSP; P01679; 2FBJ.
HSSP; P01679; 1q; 1.
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InterPro; IPR003006; Pfam; PF00047; ig; 1.
Immunoglobulin V regio
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Mammalia; Eutheria; Rodentia;
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"k Chain variable regions
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                                                                                                                                                      combining sites.";
                                                                                                                                                                Rudikoff S., Rao D.N., Glauder "Kappa Chain joining segments
                                                                                                                                                                                                                     NCBI_TaxID=10090;
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21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1996 (Rel. 38, Last annotation
IG KAPPA CHAIN V-VI REGION SAPC 10.
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                                                                                                                                                                                                SEQUENCE
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                                                                                                                                              Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                        GALACTAN
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96
106
                                                                                                                                                                pubMed=6776525;
D.N., Glaudemans oning segments and
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11627
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         FRAMEWORK 1.

COMPLEMENTARITY-DETERMINING 1

FRAMEWORK 2.

COMPLEMENTARITY-DETERMINING 2

FRAMEWORK 3.

COMPLEMENTARITY-DETERMINING 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                         8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 411.5;
Pred. No. 6.
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COMPLEMENTARITY-DETERMINING
FRAMEWORK 4.
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COMPLEMENTARITY-DETERMINING
FRAMEWORK 2.

COMPLEMENTARITY-DETERMINING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BY SIMILARITY
                                                                                                                                    77:4270-4274(1980). WAS ISOLATED FROM A
                                                                                                                                                                                                                               Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                   PRT;
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structural diversity of
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Matches 81
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Best Local S
Matches 85
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NON_TER
SEQUENCE
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DOMAIN
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                       Pfam; PF00047; Immunoglobulin
                                                                                                                                                                                                                                                                                                               modified
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                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-83271467; PubMed-6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Mil
MarkA sequences define an unusually restricted
phenyloxazolone and its early diversification.'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-AUG-1987 (Rel.
15-JUL-1999 (Rel.
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13-AUG-1987
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                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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                                                VLTQSPAIMSASPGEKVTMTCRASSSVSSSYLHWYRQKSGASPKLWIYSTSNLASGVPAR 62
FSGSGSATSYSLTITRMQAEDAATYYCQQWSSYPPMLTFGAGTKLELK
                   FSGSGSGTSYSLTISSVEAEDAATYYCQQYSGY---RTFGGGTKLEIK 107
                                     LLTQSPAIMSASPGQKVTMTCSASSSV--SYMYWYQQKPGSSPRLLIYDTSNLASGVPVR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ARFSGSGSGTSYSLTISSVEAEDAATYYCQQYS-GYRTFGGGTKLEIKR 108
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                                                                                          Similarity
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                                                                                                                                                                                                                                                                                        non-profit institutions as long as its content is in no way and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
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107
107
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                                                                                 Conservative
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V region; Hybridoma
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78.0%;
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                                                                               Score 407.5;
Pred. No. 1.3e
L2; Mismatches
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Pred.
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BY SIMILARITY.
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No. 6.1e-26;
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Matches 84
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KV6B_MOUSE
P01676;
21-JUL-1986
21-JUL-1986
15-JUL-1999
                                                                                                                               KV6C_MOUSE STAN
P01677;
21-JUL-1986 (Rel. (
21-JUL-1986 (Rel. (
15-JUL-1999 (Rel. 3)
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                                                                                      Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae
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DOMAIN
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DOMAIN
         Biochemistry 17:5555-559(1978).
                  proteins."
                            MEDLINE-7908280; PubMed-103573;
Rao D.N., Rudikoff S., Potter M.
"k Chain variable regions from t
                                                                                                                   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG KAPPA CHAIN V-VI REGION TEPC 601/TEPC 191.
                                                                                                                                                                                  MOUSE
                                                                              NCBI_TaxID=10090;
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Mammalia; Eutheria;
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Immunoglobulin V ro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biochemistry 17:5555-5559(1978)
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG KAPPA CHAIN V-VI REGION XRPC 24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR003006; -.
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                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                  ARFSGSGSTSYSLTISSVEAEDAATYYCQQYS-GYRTFGGGTKLEIKR 108
                                                                                                                                                                                                                                                                          ENVLTQSPAIMSASPGEKVTMTCRASSSVSSSYLHWYRQKSGASPKLWIYSTSNLASGVP 60
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23
33
48
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87
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11584
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Rodentia;
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77.1%;
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3; Mismatches
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BY SIMILARITY.
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Sciurognathi;
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                             galactan
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thi; Muridae; Murinae; Mus
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                                                                                         Muridae;
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                             binding
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                                                                                                                                                                                                                                                                                                                           Length
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                                                                                         Murinae;
                             myeloma
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RESULT 15
KV6E_MOUSE
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Best Local
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P01679;
21-JUL-1986
21-JUL-1986
15-JUL-1999
IG KAPPA CHAN
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                                                                  study at 2.6-A resolution proteins 1:74-80(1986).
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DOMAIN
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DOMAIN
                                     PIR; A01942; KVMSJ5.
PDB; 2FBJ; 15-OCT-90.
                                                                                                                     X-RAY CRYSTALLOGRAPHY (2.6 ANGSTROMS).
MEDLINE-88217852; PubMed-3449853;
Suh S.W., Bhat T.N., Navia M.A., Cohen
                                                                                                                                                                    MEDLINE=81054757; PubMed=6776525; Rudikoff S., Rao D.N., Glaudemans "Kappa Chain joining segments and combining sites.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE (TEPC 191).

MEDLINE-8105475; Pubmed-6776525;

Rudikoff S., Rao D.N., Glaudemans C.P.J., Potter M.;

"Kappa Chain joining segments and structural diversity of antibody combining sites.";

Proc. Natl. Acad. Sci. U.S.A. 77:4270-4274(1980).

-!- MISCELLANEOUS: THE TWO SEQUENCES ARE IDENTICAL.

-!- MISCELLANEOUS: THESE CHAINS WERE ISOLATED FROM MYELOMA PROTEINS
Immunoglobulin V re
                                                                                                            Davies D.R.;
                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoglobulin
DOMAIN 1
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Pfam; PF00047; ig; 1
Immunoglobulin V reg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                THAT BIND GALACTAN.
PIR; A01941; KVMSX4.
                             InterPro;
                                                                                       "The galactan-binding immunoglobulin study at 2.6 A resolution.";
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                                                                                                                                                               Proc. Natl.
                                                                                                                                                                                                                  SEQUENCE
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nes 83; Conserv
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-JUL-1986 (Rel. 01, Last sequence update)
-JUL-1999 (Rel. 38, Last annotation update)
KAPPA CHAIN V-VI REGION J539.
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